

Figure 1

00067638-042999

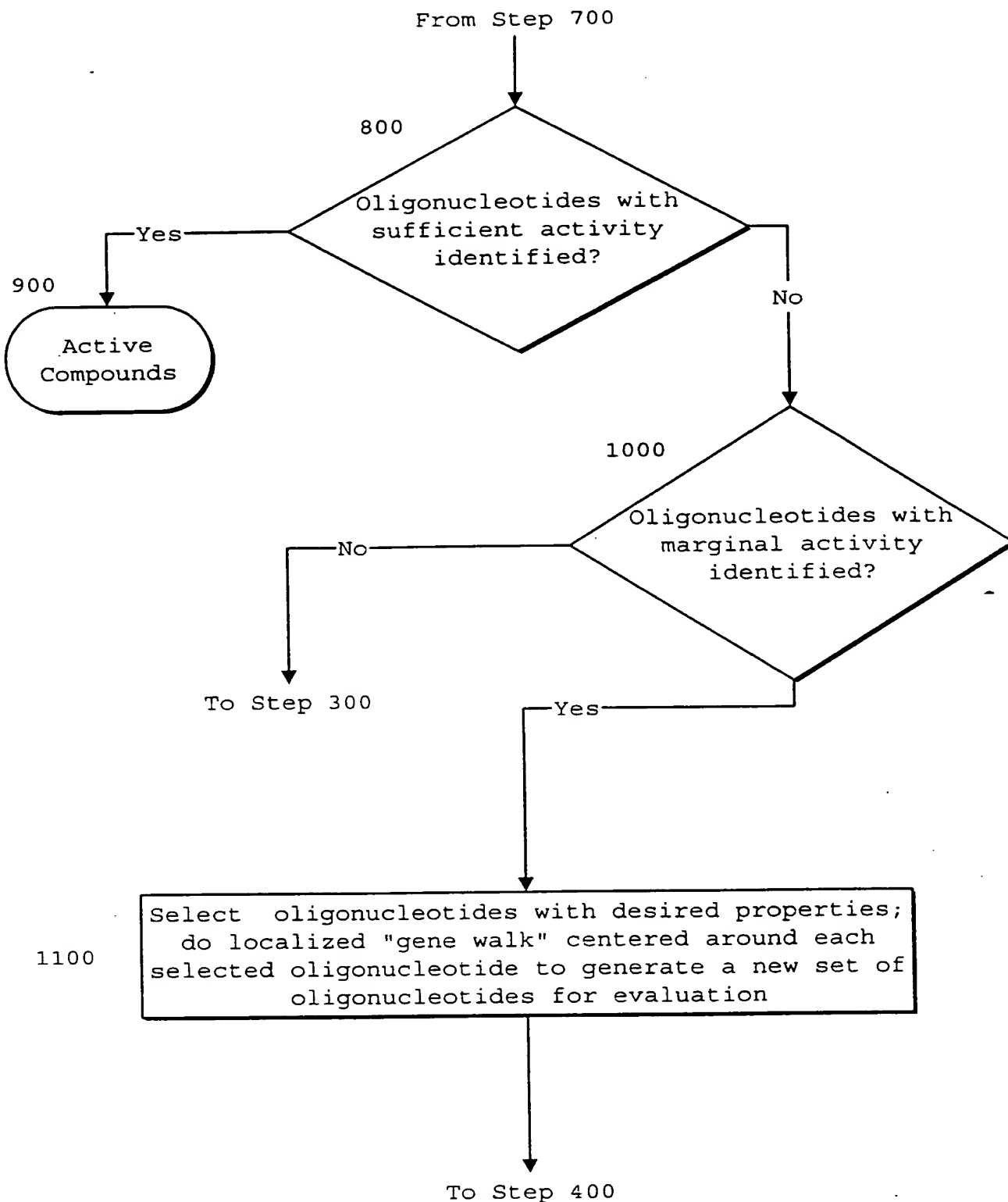


Figure 2

09067538 042898
868240" 8E9/9060

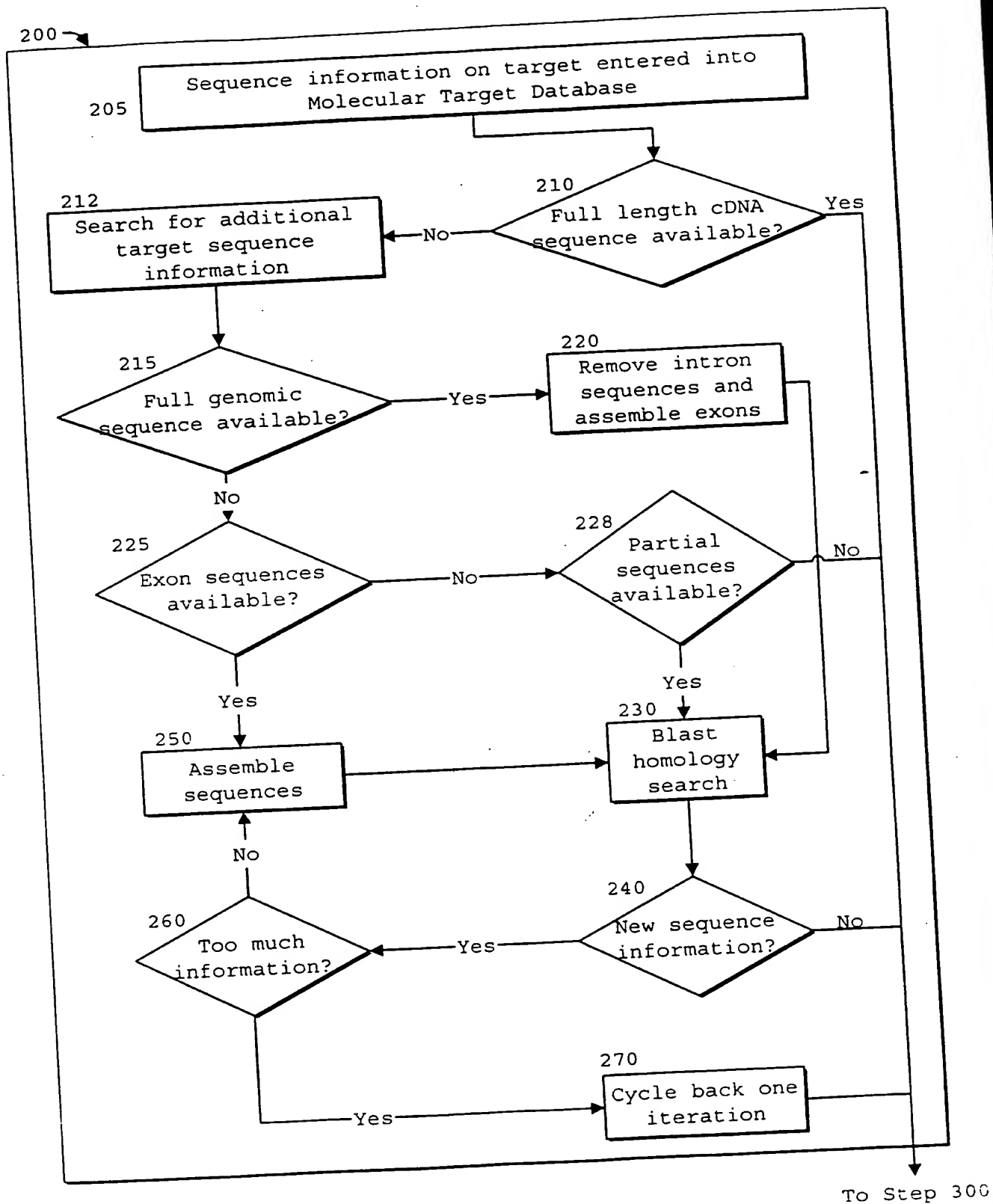


Figure 3

300

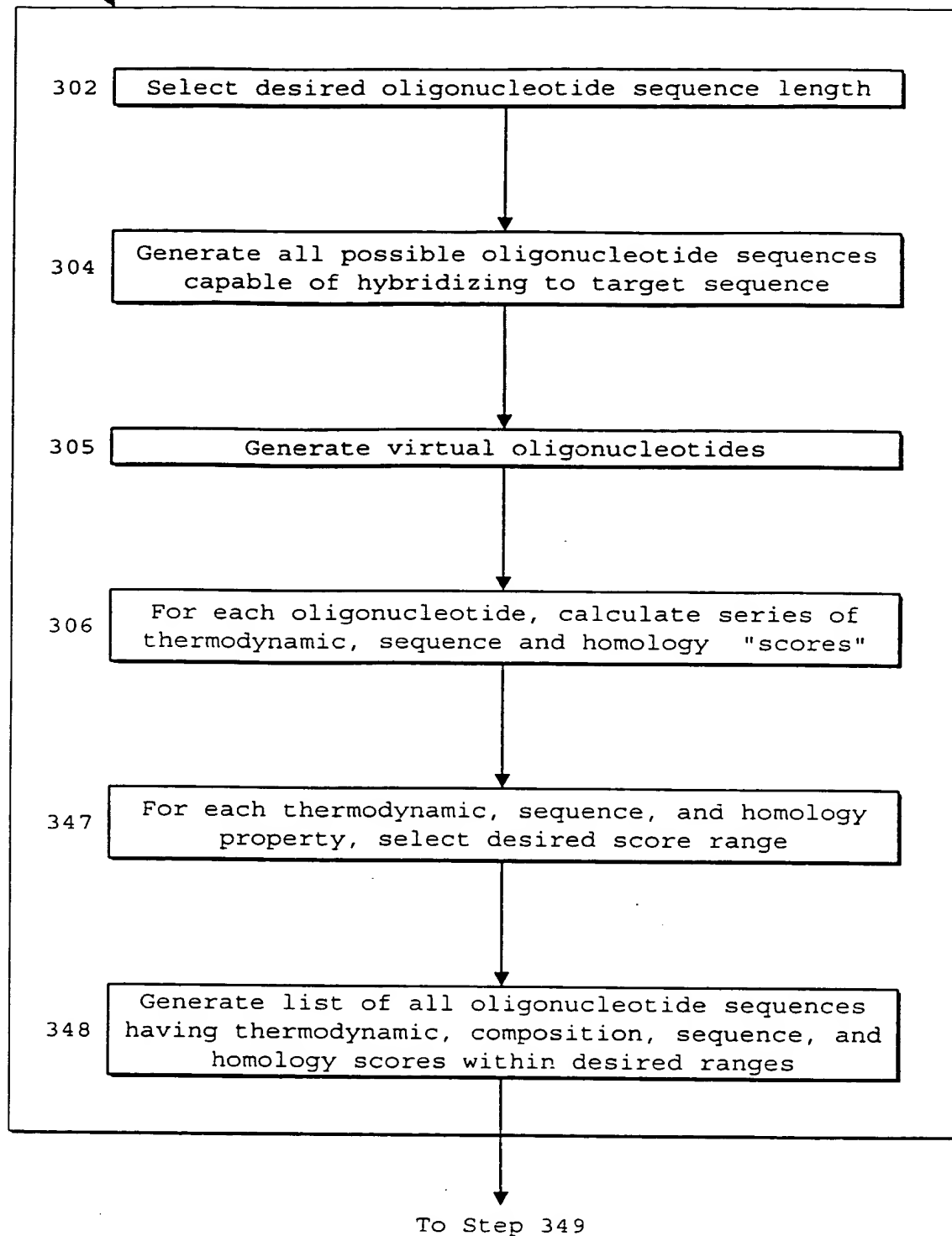


Figure 4

09067639-042899

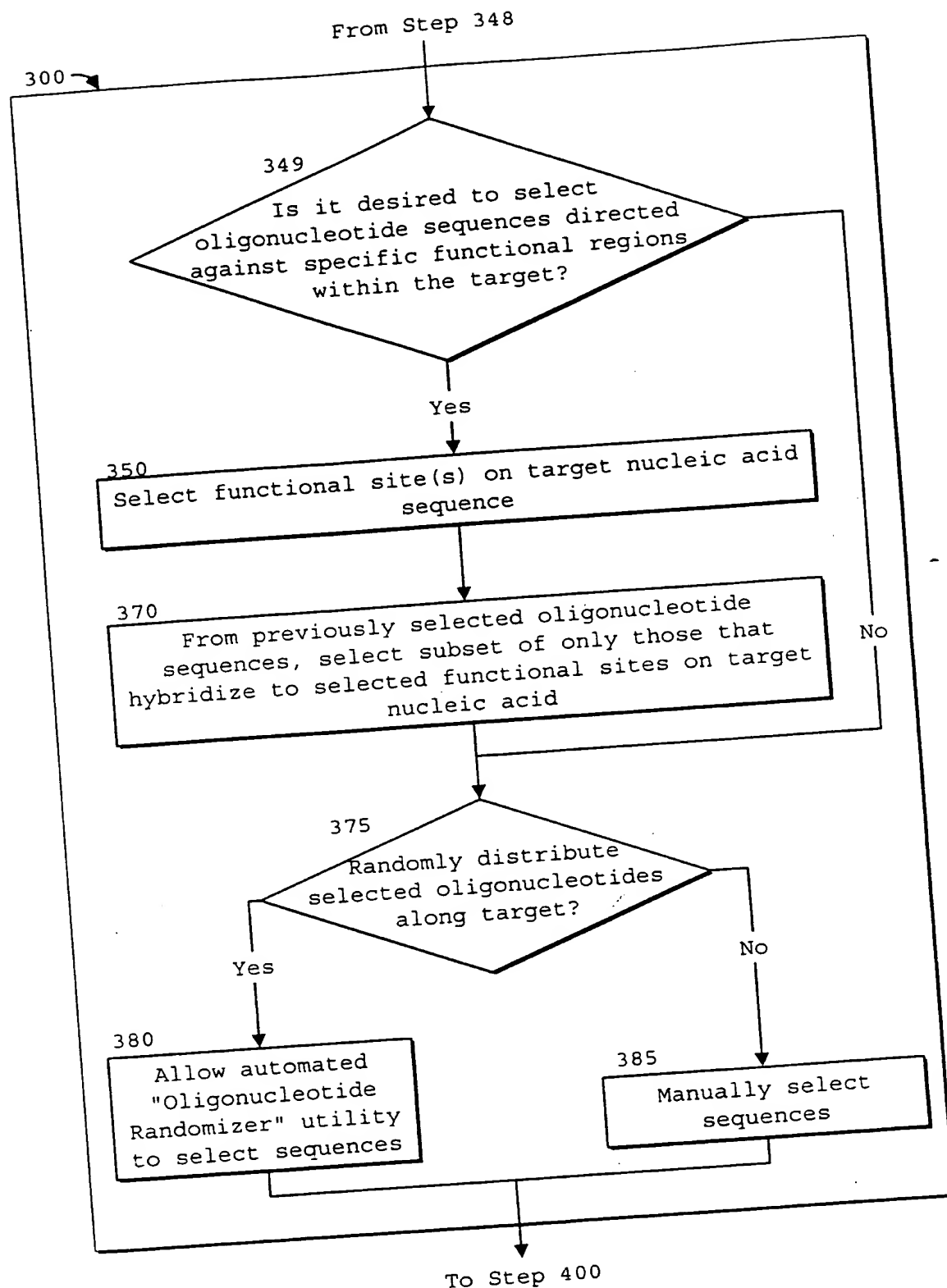


Figure 5

00067638-042898

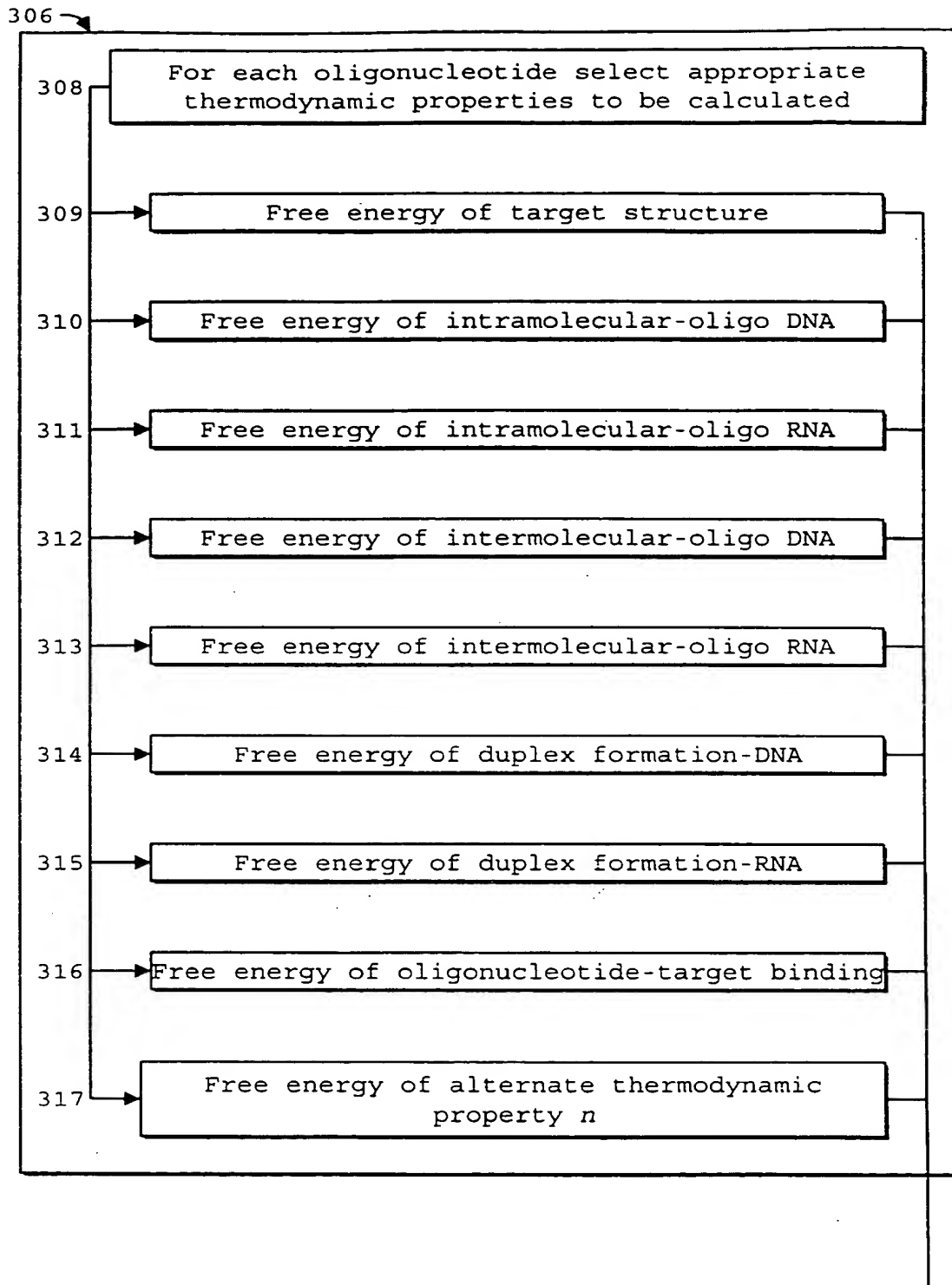
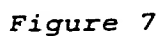
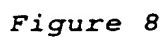


Figure 6





353	Transcription start site (5' cap)
354	5' UTR
355	Start codon
356	Coding region
357	Stop codon
358	3' UTR
359	5' splice site
360	3' splice site
361	Exon N (N = number of exon)
362	Intron N (N = number of intron)
363	mRNA stabilization signal
364	mRNA destabilization signal
365	Poly-adenylation signal
366	Poly-A addition site
367	Poly-A tail
368	Gene sequence 5' of known pre-mRNA
369	Alternate secondary structure property n

Figure 9

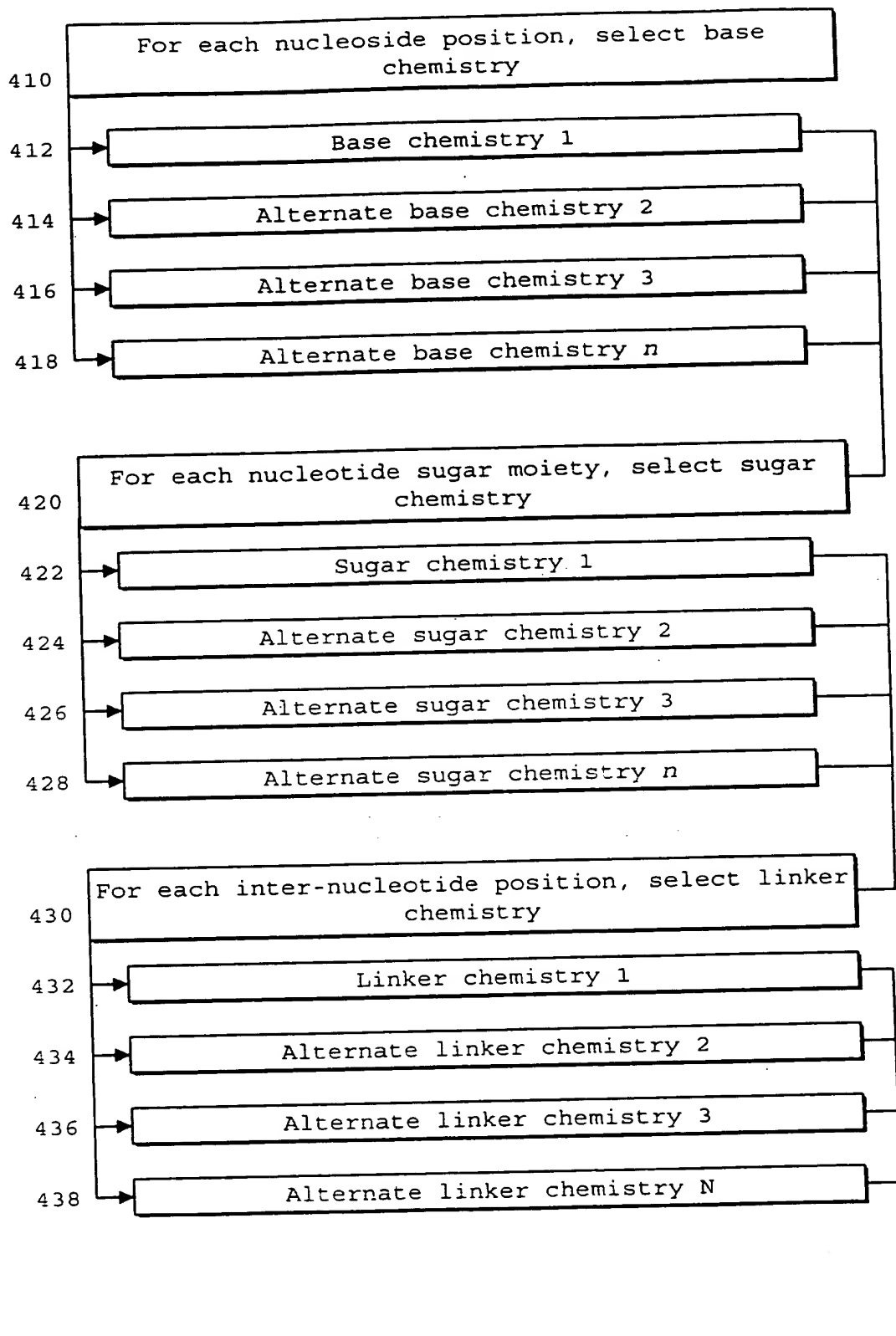


Figure 10

09067638 042899

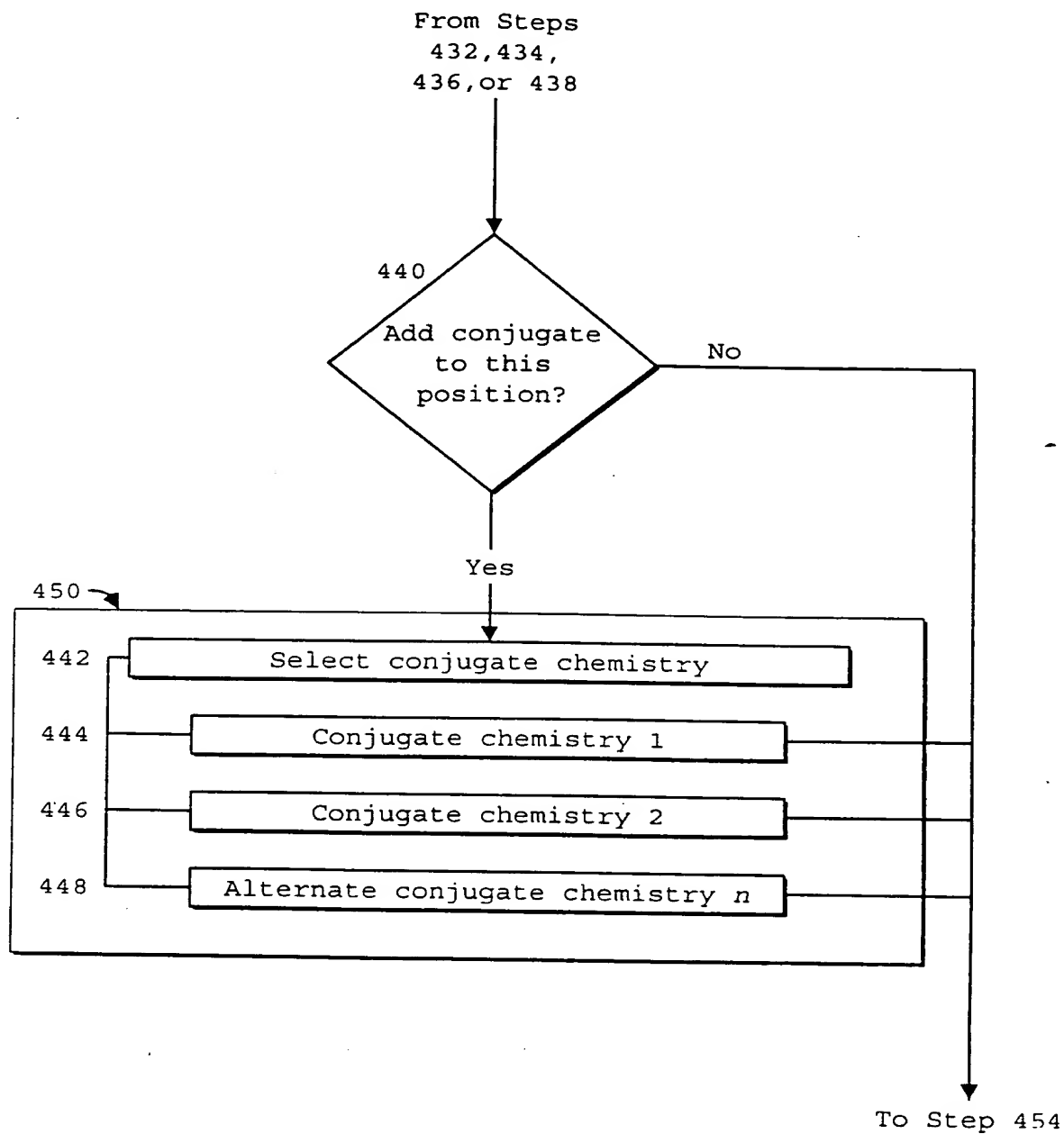


Figure 11

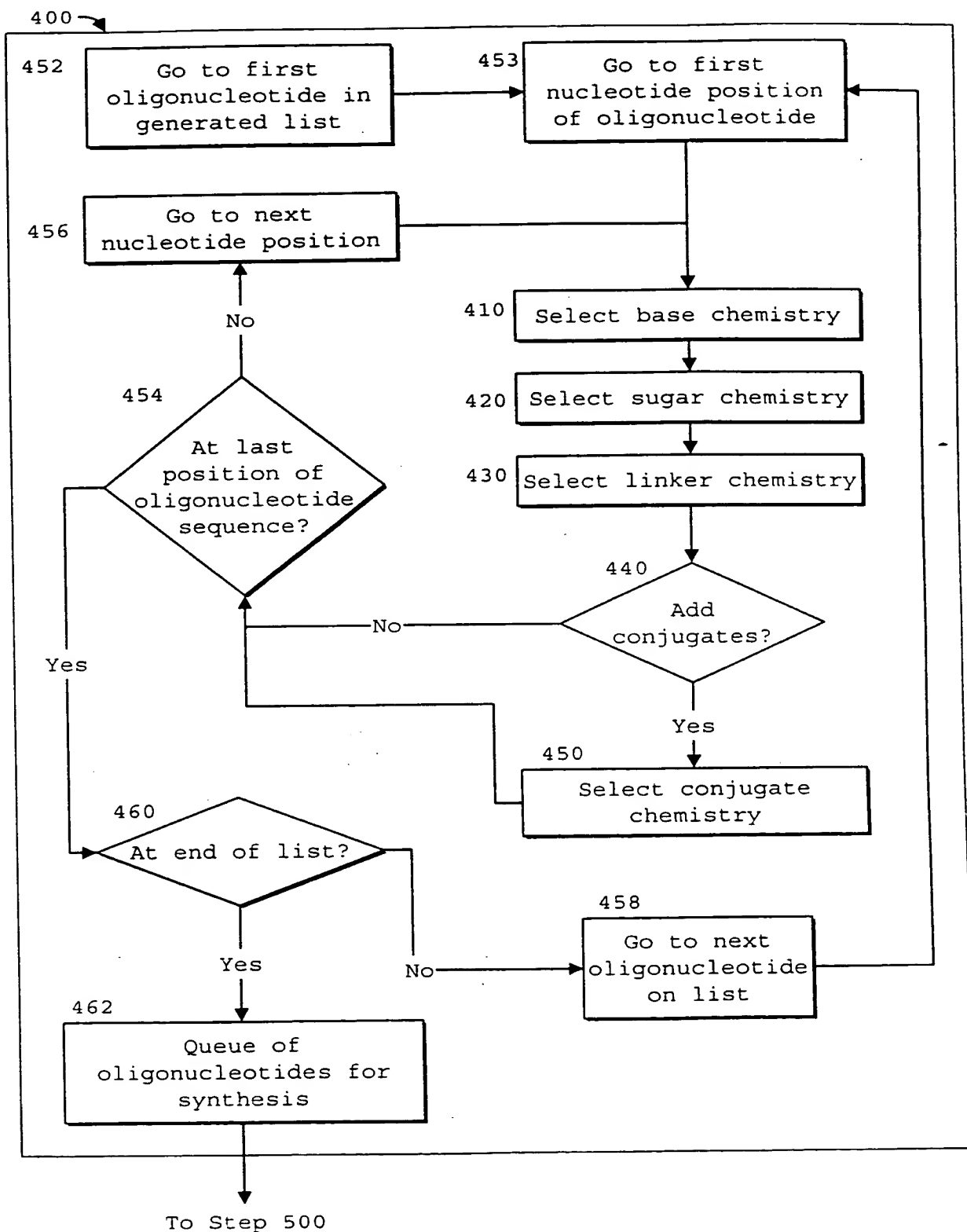


Figure 12

09067639.042898

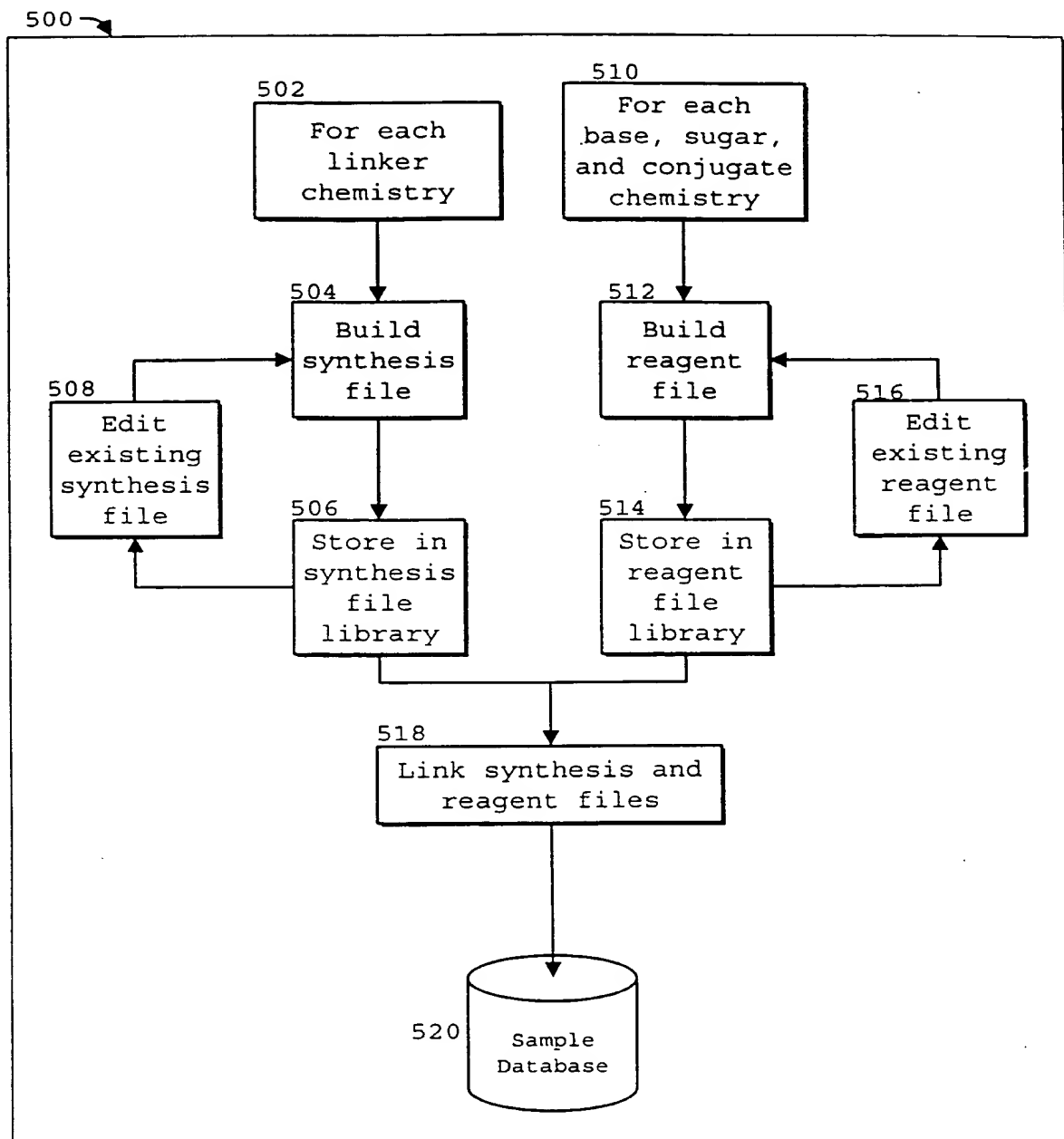


Figure 13

00067638-042890

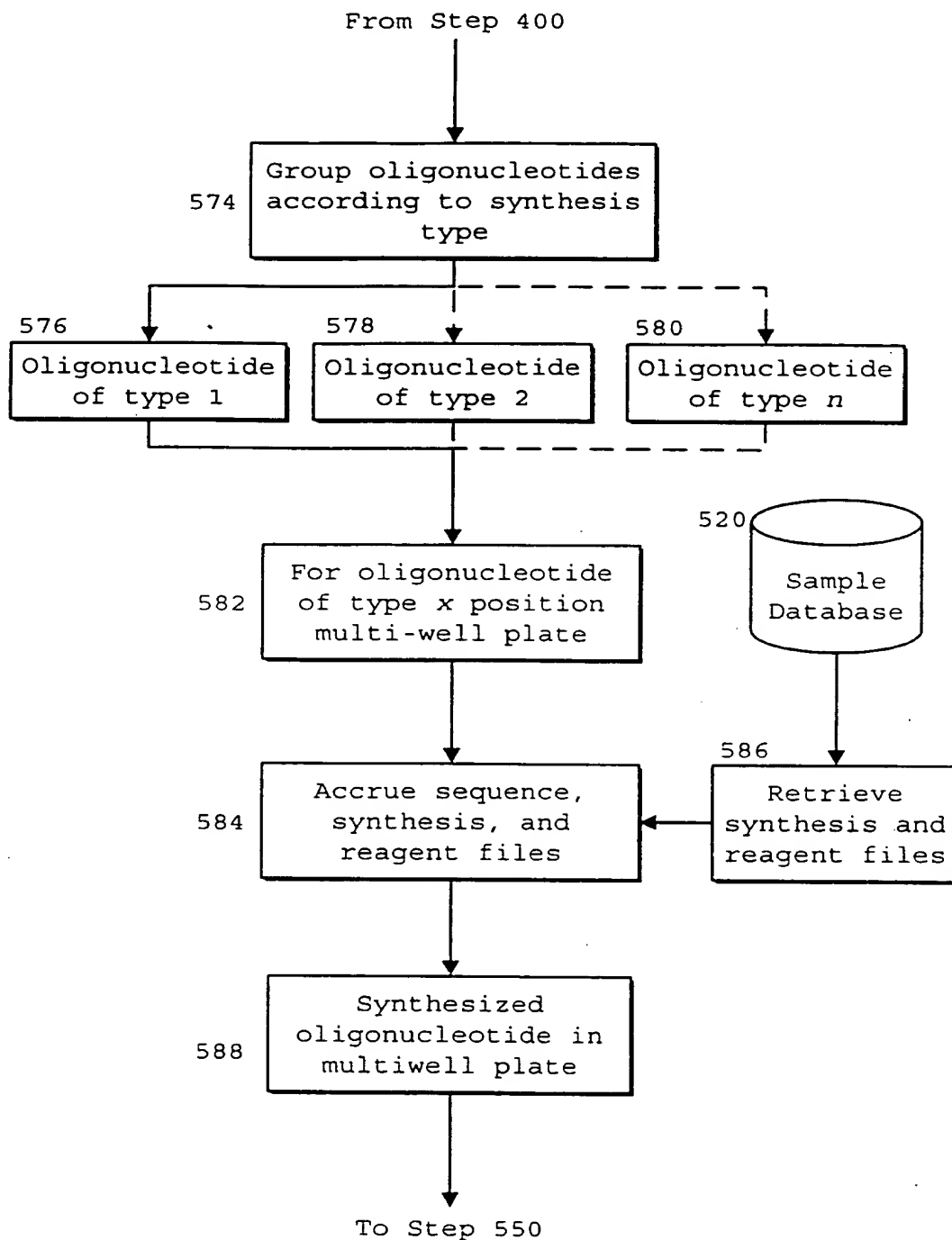


Figure 14

00067638 04290

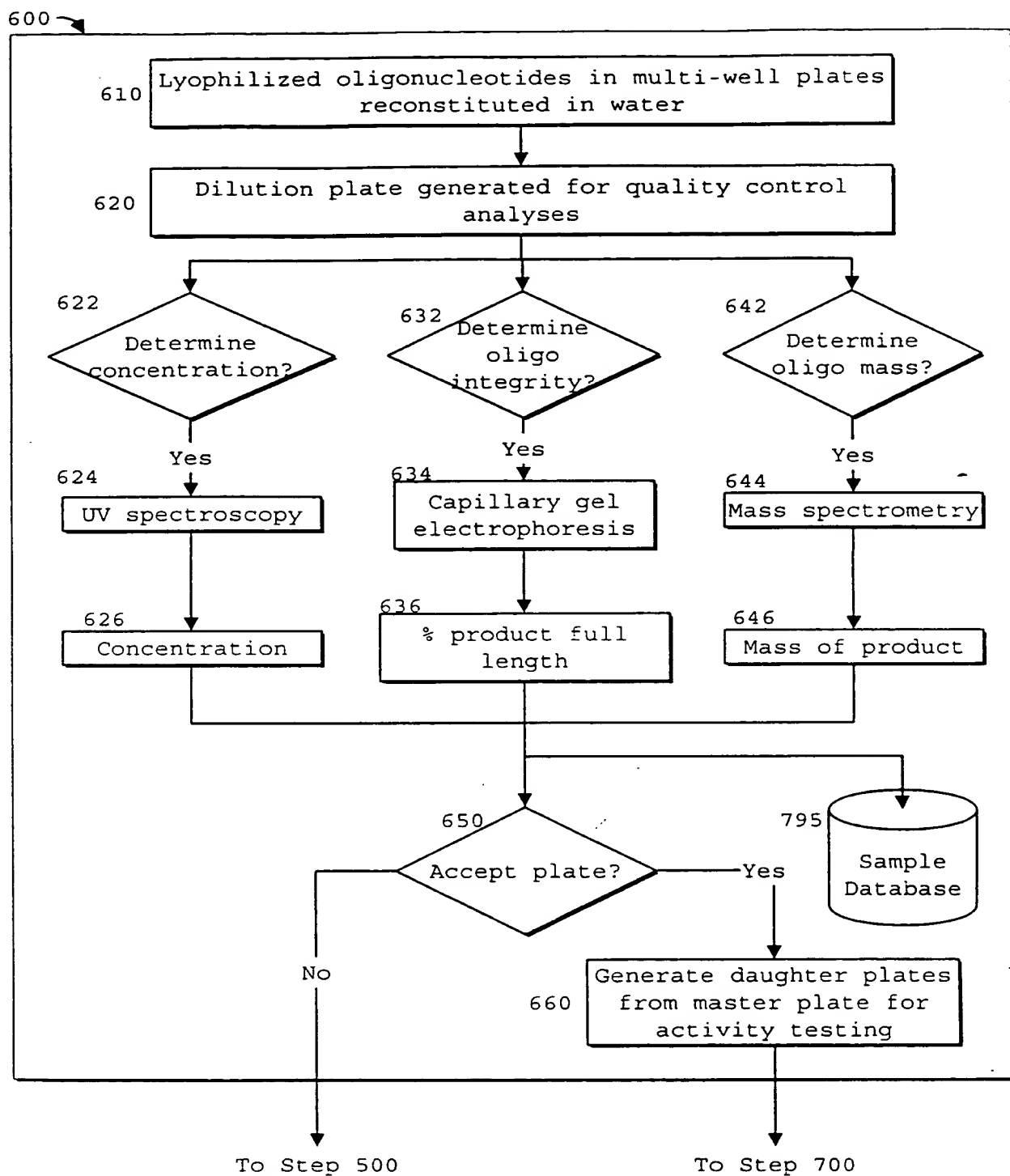


Figure 15

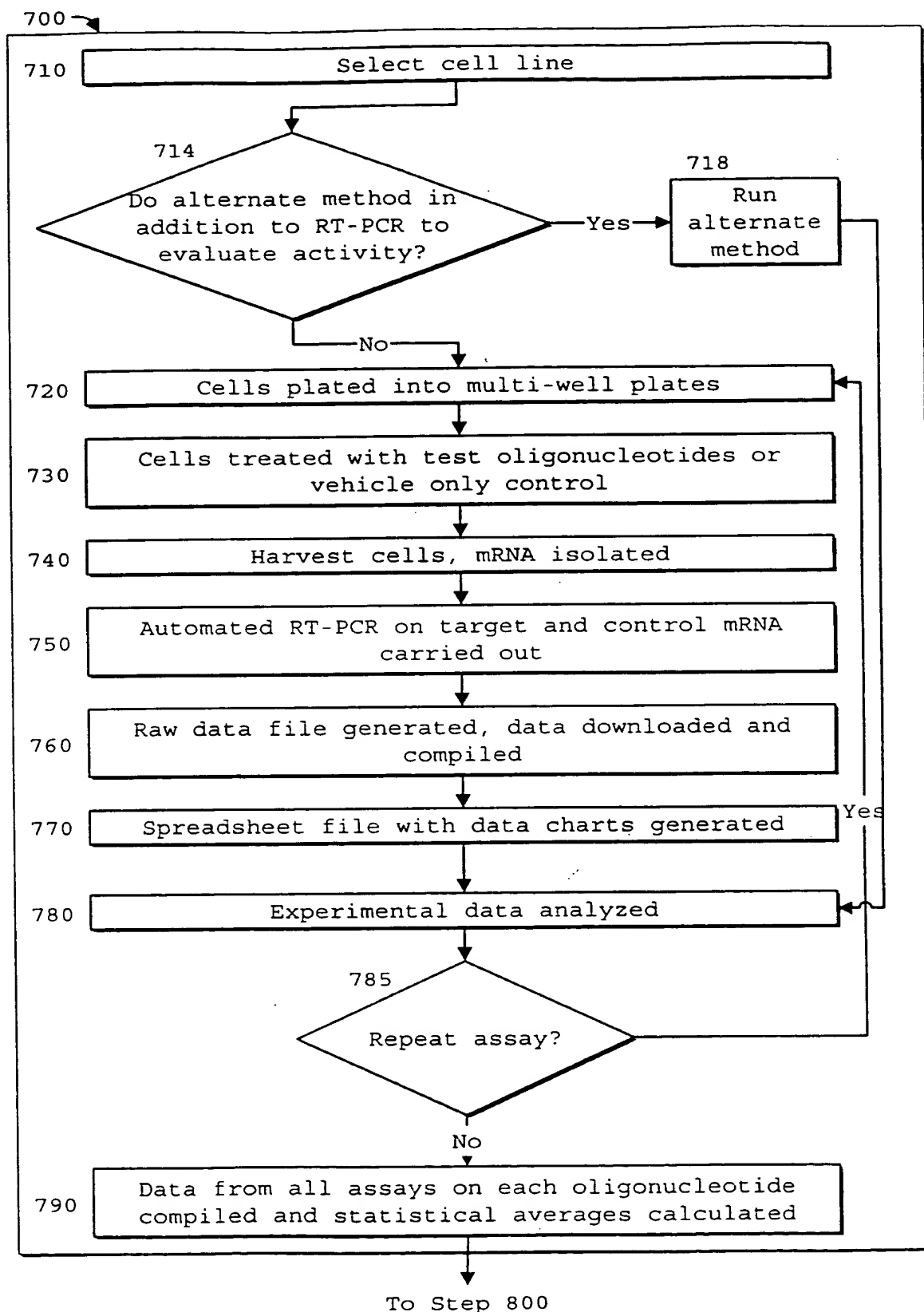


Figure 16

09067638-042899

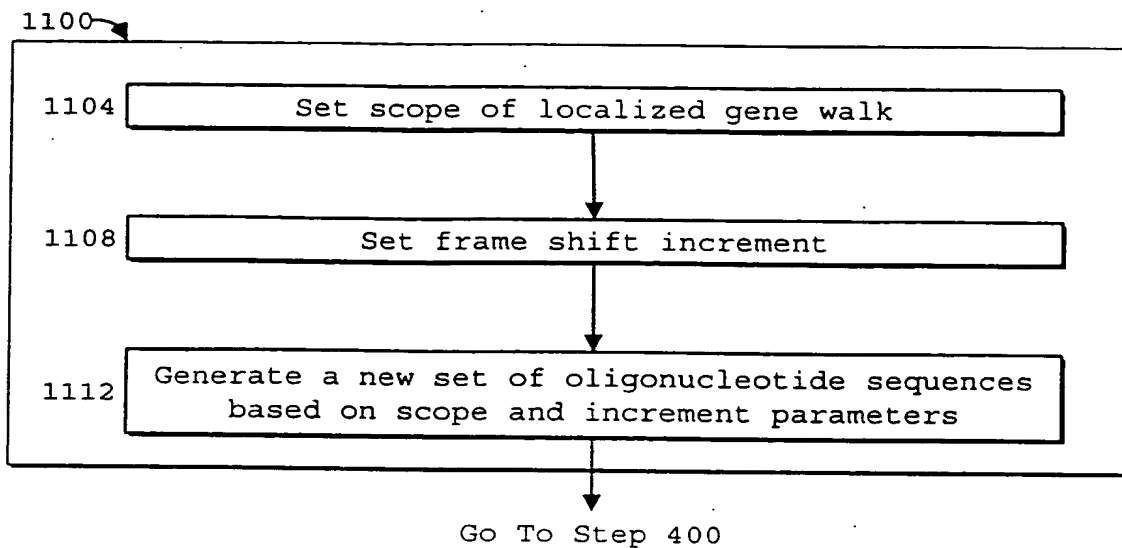


Figure 17

05067638-042898

668240" 8E929060

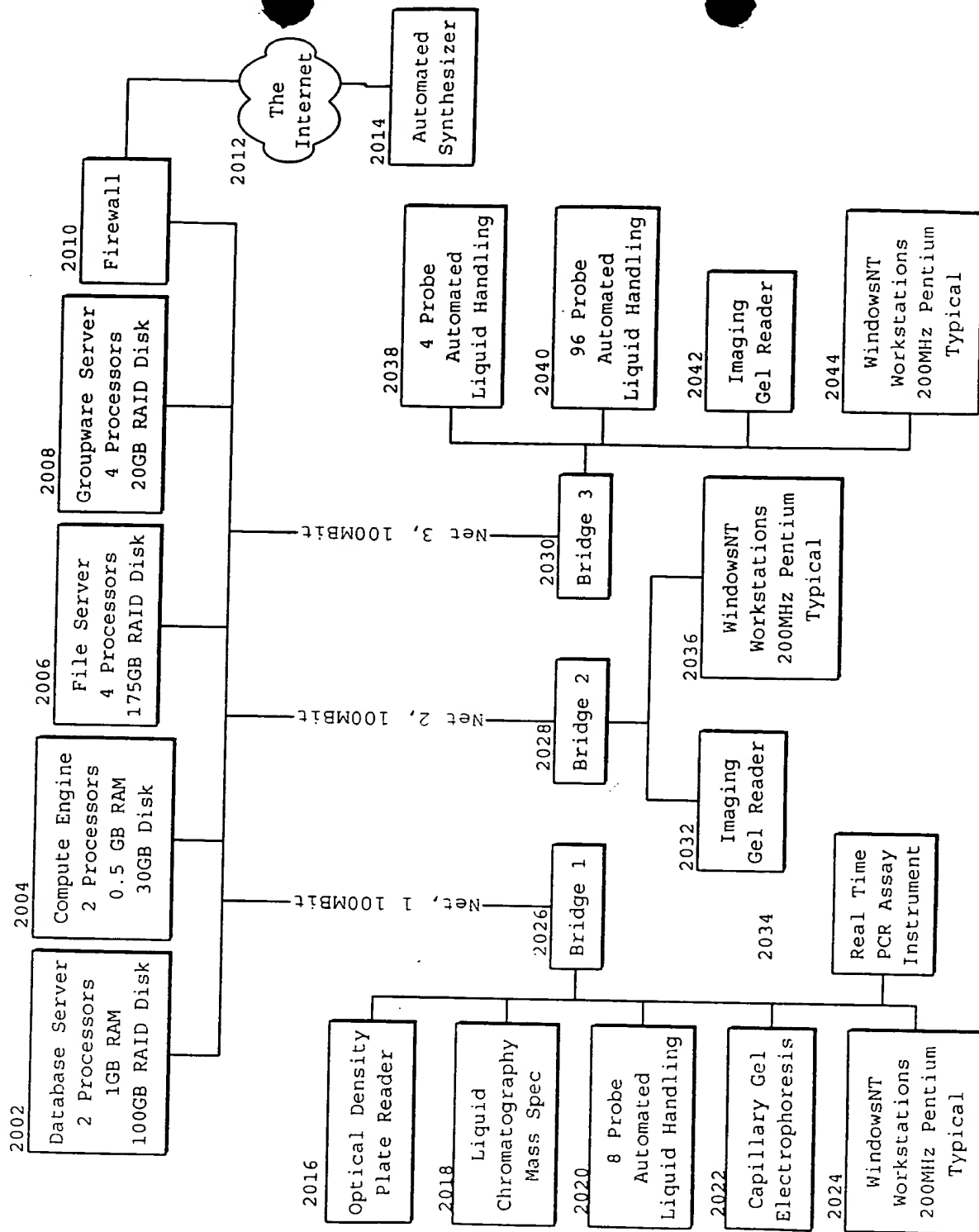


Figure 18

Figure 19

36340" 66929060

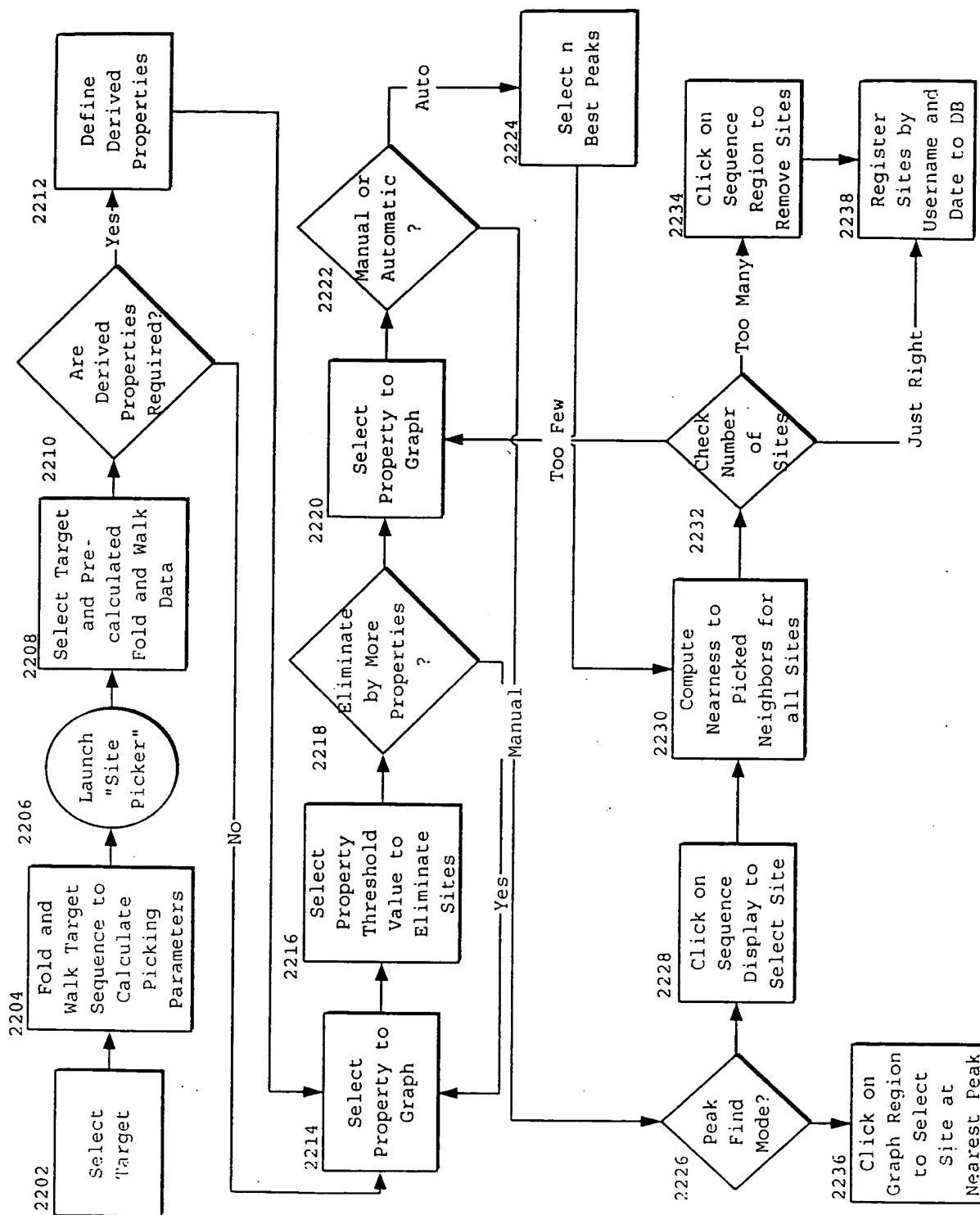
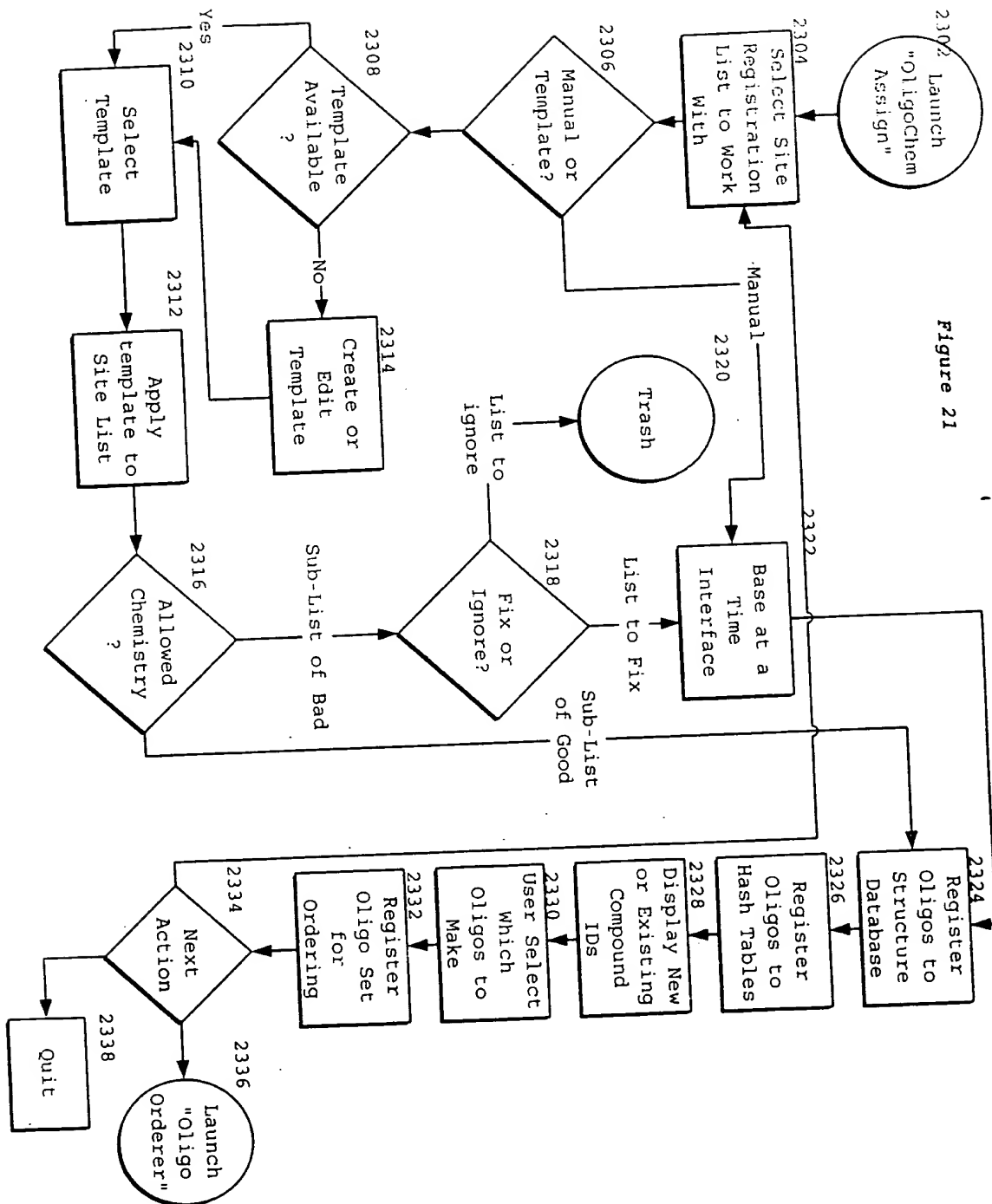


Figure 20

Figure 21



09067638 . 042398

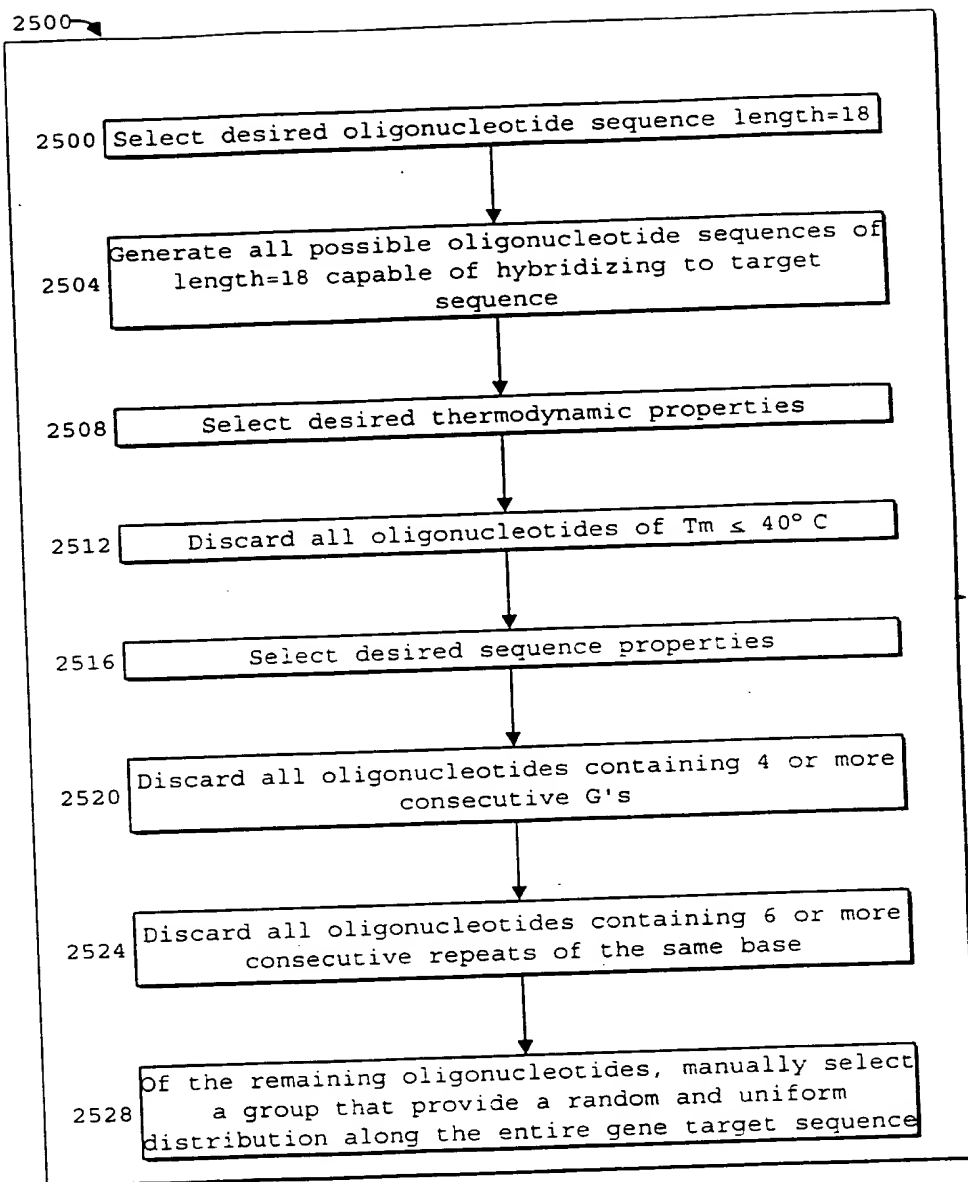


Figure 22

09067638-042888

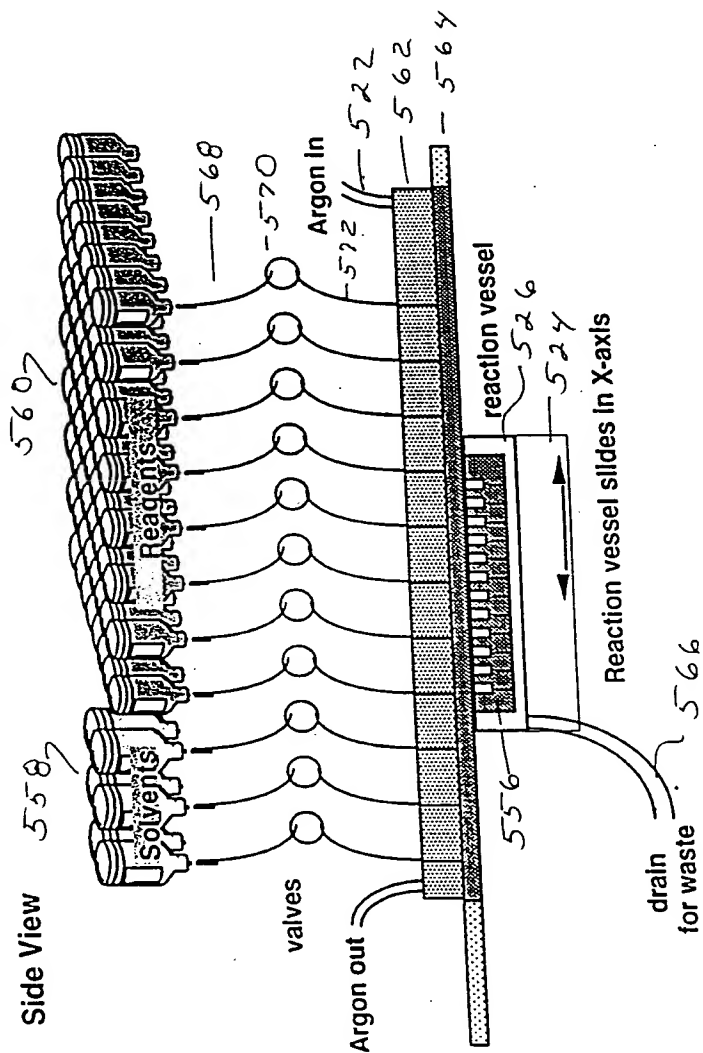


Figure 23

1306240" 8E929060

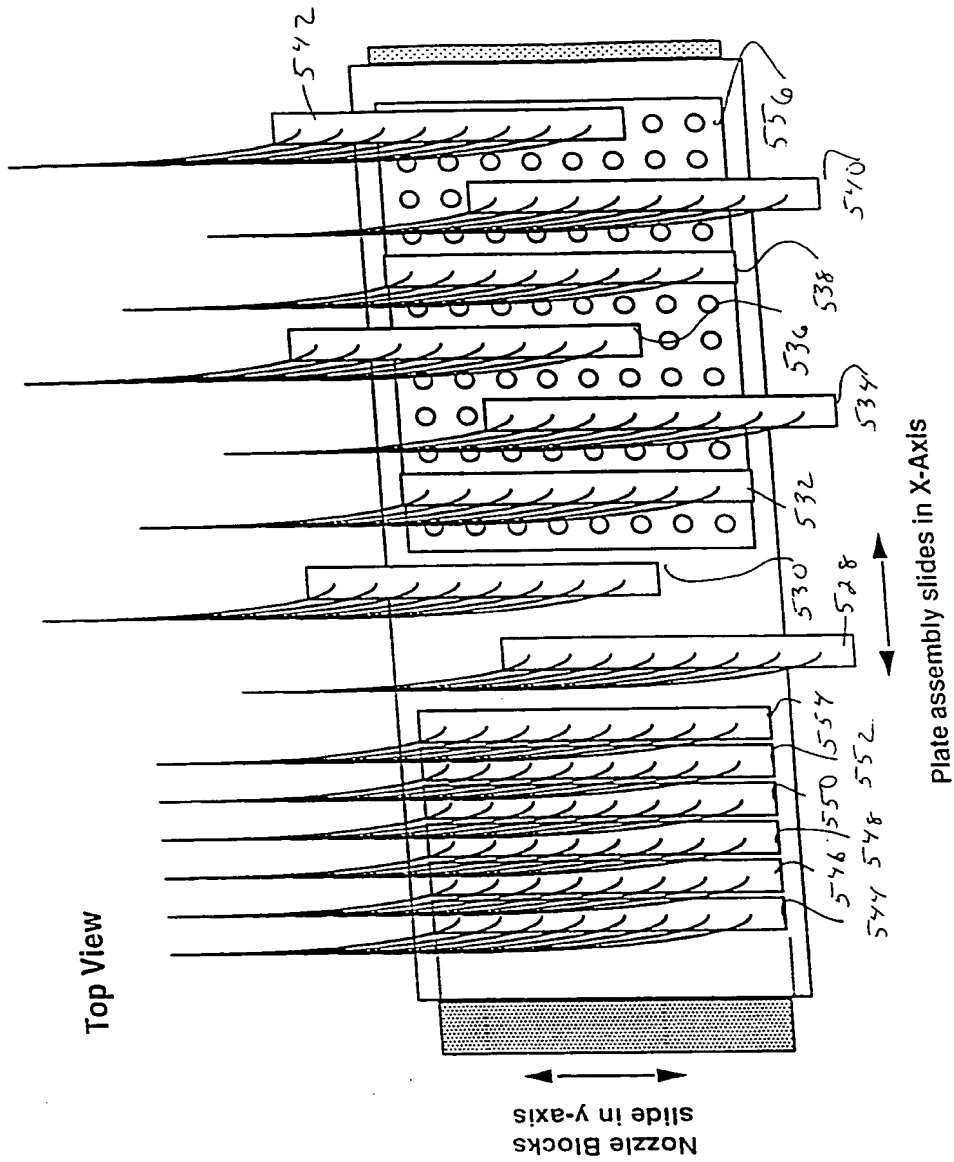


Figure 24